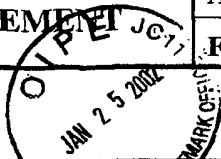


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	Applicant(s): Eric T. BALDWIN et al.	Confirmation No.: 7868
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U.S. PATENT DOCUMENTS

Examiner Initial	Document Number	Date	Name	Class	Subclass	Filing Date If Appropriate
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FOREIGN PATENT DOCUMENTS

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Examiner Initial	Document Number	Date	Country	Class	Subclass	Translation	
NE	0 879 879 A2	11/25/98	EPO			Yes	No

OTHER DOCUMENTS (Including Authors, Title, Date, Pertinent Papers, etc.)

Examiner Initial	Document Description
NE	Becker et al., "Structure of Peptide Deformylase and Identification of the Substrate Binding Site," <i>The Journal of Biological Chemistry</i> , 273(19):11413-11416 (1998).
NE	Becker et al., "Iron center, substrate recognition and mechanism of peptide deformylase," <i>Nature Structural Biology</i> , 5(12):1053-1058 (1998).
NE	Brizzard et al., "Immunoaffinity Purification of FLAG [®] Epitope-Tagged Bacterial Alkaline Phosphatase Using a Novel Monoclonal Antibody and Peptide Elution," <i>BioTechniques</i> , 16(4):730-735 (1994).
NE	Chang et al., "Methionine Aminopeptidase Gene of <i>Escherichia coli</i> Is Essential for Cell Growth," <i>Journal of Bacteriology</i> , 171(7):4071-4072 (1989).
NE	Chen et al., "Mechanistic Studies on the Aminopeptidase from <i>Aeromonas proteolytica</i> : A Two-Metal Ion Mechanism for Peptide Hydrolysis," <i>Biochemistry</i> , 36(14):4278-4286 (1997).
NE	Chiang et al., "Expression and Purification of General Transcription Factors by FLAG Epitope-Tagging and Peptide Elution," <i>Peptide Research</i> , 6(2):62-64 (1993).
NE	Dardel et al., "Solution Structure of Nickel-peptide Deformylase," <i>Journal of Molecular Biology</i> , 280(3):501-513 (1998).
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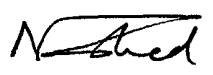
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	Groche et al., "Isolation and Crystallization of Functionally Competent <i>Escherichia coli</i> Peptide Deformylase Forms Containing either Iron or Nickel in the Active Site," <i>Biochemical and Biophysical Research Communications</i> , 246(2):342-346 (1998).
NE	Hopp et al., "A Short Polypeptide Marker Sequence Useful for Recombinant Protein Identification and Purification," <i>Biotechnology</i> , 6(10):1204-1210 (1988).
NE	Hu et al., "H-Phosphonate Derivatives as Novel Peptide Deformylase Inhibitors," <i>Bioorganic & Medicinal Chemistry Letters</i> , 8:2479-2482 (1998).
NE	Hu et al., "Determination of Substrate Specificity for Peptide Deformylase through the Screening of a Combinatorial Peptide Library," <i>Biochemistry</i> , 38(2):643-650 (1999).
NE	Laemmli, "Cleavage of Structural Proteins during the Assembly of the Head of Bacteriophage T4," <i>Nature</i> , 227(5259):680-685 (1970).
NE	Lazennec et al., "Formate Dehydrogenase-Coupled Spectrophotometric Assay of Peptide Deformylase," <i>Analytical Biochemistry</i> , 244:180-182 (1997).
NE	Meinzel et al., "Mapping of the Active Site Zinc Ligands of Peptide Deformylase," <i>Journal of Molecular Biology</i> , 254(2):175-183 (1995).
NE	Meinzel et al., "A New Subclass of the Zinc Metalloproteases Superfamily Revealed by the Solution Structure of Peptide Deformylase," <i>Journal of Molecular Biology</i> , 262(3):375-386 (1996).
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NE	Meinzel et al., "Design and Synthesis of Substrate Analogue Inhibitors of Peptide Deformylase," <i>Biochemistry</i> , 38(14):4287-4295 (1999).
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NE	QIAexpress® - The Complete System Ni-NTA Technology and the 6xHis Tag. Datasheet [online]. Qiagen [retrieved on 2001-11-06]. Retrieved from the Internet: <URL:www.qiagen.com/catalog/chapter_03/chap3.asp>, 3 pages.

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Examiner Initial	Document Description
NE JAN 2 5 2002 PATENT & TRADEMARK OFFICE	QIAexpress® Expression System. Datasheet [online]. Qiagen [retrieved 2001-11-06]. Retrieved from the Internet: <URL:www.qiagen.com/catalog/chapter_03/chap3.asp>, 5 pages.
NE	QIAexpress® Protein Purification System. Datasheet [online]. Qiagen [retrieved on 2001-11-06]. Retrieved from the Internet: <URL:www.qiagen.com/catalog/chapter_03/chap3.asp>, 5 pages.
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NE	Wei et al., "Continuous Spectrophotometric Assay of Peptide Deformylase," <i>Analytical Biochemistry</i> , 250:29-34 (1997).

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